



RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number: 09/156,580
Art Unit / Team No. : 01PE
Date Processed by STIC: 9/30/98

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,**
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY**

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

ARTI SHAH 703-308-4212

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/156,580

DATE: 09/30/98
TIME: 11:29:13

Input Set: I156580.RAW

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

Does Not Comply
Corrected Diskette Needed

new format

E--> OK

1 <110> NATIONAL INSTITUTE OF AGROBIOLOGICAL RESOURCES, MINISTRY OF
2 AGRICULTURE, FORESTRY AND FISHERIES
3 <120> A GENE FOR TRANSCRIPTION FACTOR CAPABLE OF ALTERING A CHARACTER
4 OF A PLANT AND USE THEREOF
5 <130> J198412287
6 <150> JP10-224852
7 <151> 1998-08-07
8 <160> 5
9 <170> PatentIn Ver. 2.0

ERRORED SEQUENCES FOLLOW

10 <210> 2
11 <211> 206
12 <212> PRT
13 <213> Petunia hybrida var. Mitchell
E--> 14 <400> 2<met "2">
15 Met Ala Gly Met Asp Arg Asn Ser Phe Asn Ser Lys Tyr Phe Lys Asn
E--> 16 5 delete -> @10 delete @15 15
17 Lys Ser Ile Met Ala Arg Gln Met Glu Tyr Leu Asn Asn Asn Gly
18 20 25 30
19 Asp Asn Asn Asn Asn Asn Val Thr Ser Ser Leu Arg Asp Asn Tyr
20 35 40 45
21 Gly Asn Glu Asp His Leu Leu Gly Gly Leu Phe Ser Trp Pro Pro Arg
22 50 55 60
23 Ser Tyr Thr Cys Ser Phe Cys Lys Arg Glu Phe Arg Ser Ala Gln Ala
24 65 70 75 80
25 Leu Gly Gly His Met Asn Val His Arg Arg Asp Arg Ala Ile Leu Arg
26 85 90 95
27 Gln Ser Pro Pro Arg Asp Ile Asn Arg Tyr Ser Leu Leu Asn Leu Asn
28 100 105 110
29 Leu Glu Pro Asn Pro Asn Phe Tyr Pro Ser His Asn Pro Ser Phe Ser
30 115 120 125
31 Arg Lys Phe Pro Pro Phe Glu Met Arg Lys Leu Gly Lys Gly Val Val
32 130 135 140
33 Pro Asn Asn His Leu Lys Ser Ala Arg Gly Arg Phe Gly Val Glu Lys
34 145 150 155 160
35 Ile Asp Ser Phe Met Gln Glu Lys Glu Cys Thr Thr Thr Val Ile Lys
36 165 170 175
37 Lys Ser Glu Phe Leu Arg Leu Asp Leu Gly Ile Gly Leu Ile Ser Glu
38 180 185 190
39 Ser Lys Glu Asp Leu Asp Leu Glu Leu Arg Leu Gly Ser Thr

Please align
amino
acid
numbers
directly
under
corresponding
amino
acids.
DO NOT
use TAB
codes.

PAGE: 2

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/156,580DATE: 09/30/98
TIME: 11:29:13

Input Set: I156580.RAW

40

195

200

205

09/186,580

<210> 3
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:primer
<220>
<221> modified base
<222> (6)
<223> i
<220>
<221> modified base
<222> (9)
<223> i
<220>
<221> modified base
<222> (12)
<223> i
<220>
<221> modified base
<222> (15)
<223> i
<400> 3
CARGCNYTNGGNGGNCAAY

18

Per 1.822(c)(2) of Sequence Rules, group all non-coding bases into 10's
10's

<210> 4
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<221> Description of Artificial Sequence:primer
<220>
<221> modified base
<222> (6)
<223> i
<220>
<221> modified base
<222> (9)
<223> i
<400> 4
YTNGGNGGNCAATGAAY

18

what about location 3?

<210> 5
<211> 17
<212> DNA
<213> Artificial Sequence
<220>
<221> primer
<220>
<221> modified base
<222> (3)
<223> i
<220>
<221> modified base
<222> (6)
<223> i
<220>

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<221> modified base
<222> (12)
<223> i *space*
<400> 5
ARNCKNARYTCNARRTC

Input Set: I156580.RAW

Line	Error/Warning	Original Text
10	E # of Seq. 5 Not Equal Actual 5	<160> 5
14	E Sequence IDs 0 and 2 differ	<400>
16	E Invalid/Missing Amino Acid Numbering	5 @ 10